

Amendments to the Specification

Please replace the paragraph beginning on page 4, lines 26 with the following rewritten paragraph

-- FIG. 2 is a representation of the secondary structure of the IMPDH monomer. (FIG 2a) Topology diagram of IMPDH domains. Secondary structure was assigned using the Kabsch and Sander (1983) algorithm along with visual inspection. The α helices and β strands that form the TIM barrel fold are labeled α_1 - α_8 and β_1 - β_8 . The remaining strands and helices are designated in alphanumeric order (e.g., α_A - α_L). The part of the structure not visible in the electron density maps is marked as "?????" "?". (FIG 2b) Stereoview ribbon diagram arranged approximately perpendicular to the axis of the TIM barrel fold; IMP is shown as a ball and stick model. --